



3/20

# FIG. 5

(SEQ ID NO: 4)

## Human Luteinizing Hormone (LH)

1                   8                   L1                   33  
SREPLRPWCHPINAILAVEKEGCPVCITVNTTICAGYCPTMMRVLQAVLP  
51                   58                   L3                   87  
PLPQVVCTYRDVRFESIRLPGCPRGVDPVVSFPVALSCRCGPCRRSTSDC  
101                   GGPKDHPLTCDHPQLSGLLFL

# FIG. 6

(SEQ ID NO: 5)

## Human Follicle Stimulating Hormone (FSH)

1                   4                   L1                   27  
NSCELTNITIAIEKEECRFCISINTTWCAGYCYTRDLVYKDPARPKitCT  
51                                   65                   L3                   81  
FKELVYETVRVPGCAHHADSLYTPVATQCHCGKCDSDSTDCTVRGLGPS  
101                   YCSFGEMKE

# FIG. 7

(SEQ ID NO: 6)

## Human Platelet-Derived Growth Factor-A (PDGF A-Chain)

1                   11                   L1                   36  
SIEEAVPAVCKTRTVIYEIPRSQVDPTSANFLIWPPCVEVKRCTGCCNTS  
51                   58                   L3                   88  
SVKCQPSRVHHRSVKVAKVEYVRKKPKLKEVQVRLEEHLECACATTSLNP  
101                   DYREEDTGRPRESGKKRKRRLKPT

*FIG. 8*  
(SEQ ID NO: 7)

17 L1 42

1 SLGSLTIAEPAMIAECKTRTEVFEISRRLIDRTNANFLVWPPCVEVQRCS

64 L3 94

51 GCCNNRNVQCRPTQVQLRPVQVRKIEIVRKKPIFKKATVTLLEDHLACKCE

101 TVAAARPVTRSPGGSQEQRAKTPQTRVTIRTVRVRRPPKGKHRKFKHTHD

151 KTALKETLGA

FIG. 9  
(SEQ ID NO: 8)

27                      L1                      50

1     APMAEGGGQNHHEVVKFMDVYQRSYCHPIETLVDIFQEYPDEIEYIFKPS

73                      L3                      99

51    CVPLMRCGGCCNDEGLECVPTESNITMQIMRIKPHQGQHIGEMSFLQHN

101   KCECRPKKDRAEQEKSVRGKGKGQKRKRKKSRYKSWSPCGPCSEERRKH

151   LFVQDPQTCKCSCKNTDSRCKARQLELNERTCRCDKPRR

FIG. 10

FIG. 11

1 HSDPARRGELSVCDSEI SEWVTAADKKTAVDMSGGTVTTVLEKVS PVKGQLK  
 51 QYFYETKCNPMGYTKEGCRGIDKRWNSQCRTTQSYVRAMLTD SKKRIGW  
 101 RFIRIDTSCVCILTIKRGR

6/20

## FIG. 12

(SEQ ID NO: 11)

### Human Neurotrophin (NT)-3

1 YAEHKSHRGEYSVCDSESLWVTDKSSAIDIRGHQVTVLGEIGKTNSPVKQ  
51 YFYETRCKEARPVKNGCRGIDDRHWN SQCKTSQTYVRASLTENNKLVGWR  
101 WIRIDTSCVCALSRKIGRT

The diagram shows the sequence of Human Neurotrophin (NT)-3. It is divided into three domains: L1 (residues 1-15), L2 (residues 16-56), and L3 (residues 57-80). The sequence is shown in three lines: 1-15, 16-56, and 57-107. The L1 domain is marked with a bracket from residue 1 to 15. The L2 domain is marked with a bracket from residue 16 to 56. The L3 domain is marked with a bracket from residue 57 to 80. The sequence ends at residue 107 with the residue WIRIDTSCVCALSRKIGRT.

## FIG. 13

(SEQ ID NO: 12)

### Human Neurotrophin (NT)-4

1 GVSETAPASRRGELAVCDAVSGWVTD RRTAVDLRGREVEVLGEVPAAGGS  
51 PLRQYFFETRCKADNAEEGGPGAGGGGCRGVDRRHVWSECKAKQSYVRAL  
101 TADAQGRVGWRWIRIDTACVCTLLSRTGRA

The diagram shows the sequence of Human Neurotrophin (NT)-4. It is divided into three domains: L1 (residues 1-18), L2 (residues 19-60), and L3 (residues 61-91). The sequence is shown in three lines: 1-18, 19-60, and 61-118. The L1 domain is marked with a bracket from residue 1 to 18. The L2 domain is marked with a bracket from residue 19 to 60. The L3 domain is marked with a bracket from residue 61 to 91. The sequence ends at residue 118 with the residue TADAQGRVGWRWIRIDTACVCTLLSRTGRA.

## FIG. 14

(SEQ ID NO: 13)

### Human Transforming Growth Factor (TGF)-β1

1 ALDTNYCFSSTEKNCCVRQLYIDFRKDLGWKWIHEPKGYHANFCLGPCPY  
51 IWSLDTQYSKVLALYNQHNP GASAAPCCVPQALEPLPIVYYVGRKPKVEQ  
101 LSNMIVRSCKCS

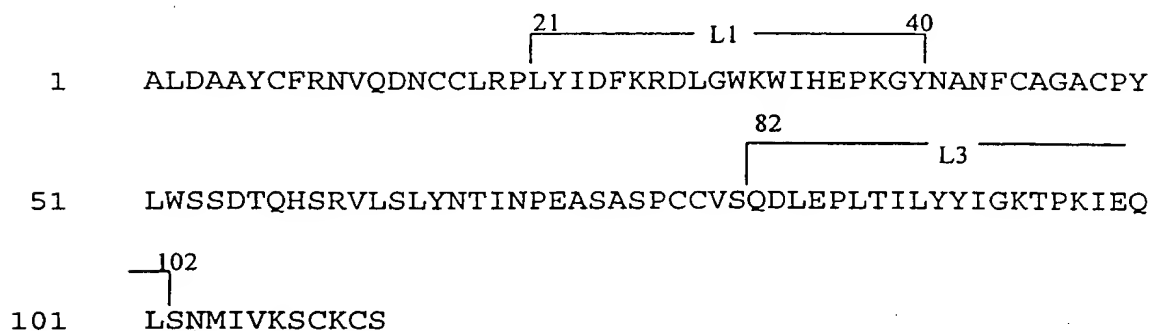
The diagram shows the sequence of Human Transforming Growth Factor (TGF)-β1. It is divided into two domains: L1 (residues 1-21) and L3 (residues 22-40). The sequence is shown in three lines: 1-21, 22-82, and 83-102. The L1 domain is marked with a bracket from residue 1 to 21. The L3 domain is marked with a bracket from residue 22 to 40. The sequence ends at residue 102 with the residue LSNMIVRSCKCS.

7/20

## FIG. 15

(SEQ ID NO: 14)

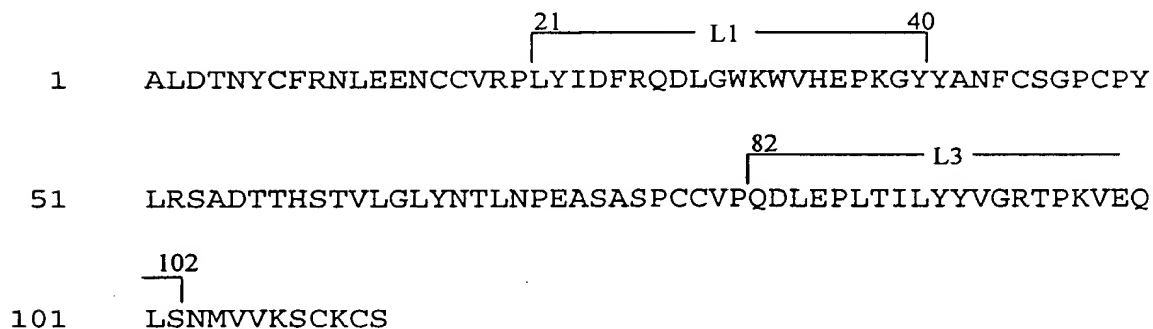
### Human Transforming Growth Factor (TGF)- $\beta$ 2



## FIG. 16

(SEQ ID NO: 15)

### Human Transforming Growth Factor (TGF)- $\beta$ 3



8/20

*FIG. 17*

(SEQ ID NO: 16)

**Human Transforming Growth Factor (TGF)- $\beta$ 4**

1 MWPLWLCWAL WVLPLAGPGA ALTEEQLLAS LLRQLQLSEV PVLDRADMEK  
51 LVIPAHVRAQ YVLLRRDGD RSRGKRFSQS FREVAGRFLA SEASTHLLVF  
101 GMEQRLPPNS ELVQAVLRLF QEPVPQGALH RHGRLSPAAP KARVTVEWLV  
151 RDDGSNRTSL IDSRLVSVHE SGWKAFDVTE AVNFWQQLSR PPEPLLQVVS  
201 VQREHLGPLA SGAHKLVRFA SQGAPAGLGE PQLELHTLDL RYGAQGD CD  
251 PEAPMTEGTR CCRQEMYIDL QGMKWAKNWV LEPPGFLAYE CVGTCQOPPE  
301 ALAFNWPFLG PRQCIASETA SLPMIVSIKE GGRTRPOVVS LPNMRVQKCS  
351 CASDGALVPR RLQHRPWCIH

*FIG. 18*

(SEQ ID NO: 17)

**Human Neurturin**

1 MQRWKAAALA SVLCSSVLSI WMCREGLLS HRLGPALVPL HRLPRTL D AR  
51 IARLAQYRAL LQGAPDAMEL RELTPWAGRP PGPRRRAGPR RRRARARLGA  
101 RPCGLRELEV RVSELGLGYA SDETVLFRYC AGACEAAARV YDLGLRRLRQ  
151 RRRLRRERVR AQPCCRPTAY EDEVSF LDAH SRYHTVHEL S ARECACV

FIG. 17



9/20

# FIG. 19

(SEQ ID NO: 18)

## Human Inhibin $\alpha$ (Common to Inhibin A and Inhibin B)

1 MVLHLLLFLL LTPQGGHSCQ GLELARELVL AKVRALFLDA LGPPAVTREG  
51 GDPGVRRLLPR RHALGGFTHR GSEPEEEEDV SQAILFPATD ASCEDKSAAR  
101 GLAQEAEEGL FRYMFRPSQH TRSRQVTSQAQ LWFHTGLDRQ GTAASNSSEP  
151 LLGLLALSPG GPVAVPMSLG HAPPHWAVLH LATSALSLLT HPVLVLLLRRC  
201 PLCTCSARPE ATPFLVAHTR TRPPSGGERA RRSTPLMSWP WSPSALRLLQ  
251 RPPEEPAAHA NCHRVALNIS FQELGWERWI VYPPSFIFHY CHGGCGLHIP  
301 PNLSLPVPGA PPTPAQPYSL LPGAQPCCAA LPGTMRPLHV RTTSDGGYSF  
351 KYETVPNLLT QHCACI

# FIG. 20

(SEQ ID NO: 19)

## Human Inhibin A - $\beta$ Subunit ( $\alpha$ - $\beta$ A Heterodimer)

1 MPLLWLRGFL LASCWIIIRS SPTPGSEGHS AAPDCPSCAL AALPKDVPNS  
51 QPEMVEAVKK HILNMLHLKK RPDVTQPVPK AALLNAIRKL HVGKVGGENGY  
101 VEIEDDIGRR AEMNELMEQT SEIITFAESG TARKTLHFEI SKEGSDLSV  
151 ERAEVWLFLK VPKANRTRTK VTIRLFQQQK HPQGSLDTGE EAEVGLKGE  
201 RSELLLSEKV VDARKSTWHV FPVSSSIQRL LDQ GKSSLDV RIACEQCQES  
251 GASLVLLGKK KKKEEEGEGK KKG GEGGAG ADEEKEQSHR PFLMLQARQS  
301 EDHPHRRRRR GLECDGKVINI CCKKQFFVSF KDIGWNDWII APSGYHANYC  
351 EGECPSHIAG TSGSSLSFHS TVINHYMRG HSPFANLKSC CVPTKLRPMS  
401 MLYYDDGONI IKKDIQNMIV EECGCS

FIG. 19

10/20

## FIG. 21

(SEQ ID NO: 20)

### Human Inhibin B - $\beta$ Subunit ( $\alpha$ - $\beta$ B Heterodimer)

1 MDGLPGRALG AACLLLLAAG WLGPEAWGSP TPPPTPAAPP P PPPPGSPGG  
51 SQDTCTSCGG FRRPEELGRV DGDFLEAVKR HILSRLQMRG RPNITHAVPK  
101 AAMVTALRKL HAGKVREDGR VEIPHLDGHA SPGADGQERV SEIISFAETD  
151 GLASSRVRLY FFISNEGNQN LFVVQASLWL YLKLLPYVLE KGSRRKVRVK  
201 VYFQEQGHGD RWNMVEKRVD LKRSGWHTFP LTEAIQALFE RGERRLNLDV  
251 QCDSQCQLAV VPVFVDPGEE SHRPFVVVQA RLGDSRHRIR KRGLECDGRT  
301 NLCCRQOFFI DFRLIGWNDW IIAPTGY YGN YCEGSCPAYL AGVPGSASSF  
351 HTAVVNQYRM RGLNPGTVNS CCIPTKLSTM SMLYFDDEYN IVKRDVPNMI  
401 VEECGCA

## FIG. 22

(SEQ ID NO: 21)

### Human Activin A ( $\beta$ A Homodimer)

1 MPLLWLRGFL LASCWIIVRS SPTPGSEGHs AAPDCPSCAL AALPKDVPNS  
51 QPEMVEAVKK HILNMLHLKK RPDVTQPVPK AALLNAIRKL HVGKVGGENGY  
101 VEIEDDIGRR AEMNELMEQT SEIITFAESG TARKTLHFEI SKEGSDLSV  
151 ERAEVWLFLK VPKANRTRTK VTIRLFQQQK HPQGS LDTGE EAEEVGLKGE  
201 RSELLLSEKV VDARKSTWHV FVSSSIQRL LDQ GKSSLDV RIACEQCQES  
251 GASLVLLGKK KKKEEEGEGK KKG GEGGAG ADEEKEQSHR PFLMLQARQS  
301 EDHPHRRRRR GLECDGKVNI CCKKQFFVSF KDIGWNDWII APSGYHANYC  
351 EGECPSHIAG TSGSSLSFHS TVINHYMRG HSPFANLKSC CVPTKLRPMS  
401 MLYYDDGQNI IKKDIONMIV EECGCS

"BEEB" 0000

11/20

## FIG. 23

(SEQ ID NO: 22)

### Human Activin B ( $\beta$ B Homodimer)

1 MDGLPGRALG AACLLLLAAG WLGPEAWGSP TPPPTPAAPP PPPPPGSPGG  
51 SQDTCTSCGG FRRPEELGRV DGDFLEAVKR HILSRLQMRG RPNITHAVPK  
101 AAMVTALRKL HAGKVREDGR VEIPHLDGHA SPGADGQERV SEIISFAETD  
151 GLASSRVRLY FFISNEGNQN LFVVQASLWL YLKLLPYVLE KGSRRKVRVK  
201 VYFQEQGHGD RWNMVEKRVD LKRSGWHTFP LTEAIQALFE RGERRLNLDV  
251 QCDSCQELAV VPVFVDPGEE SHRPFVVVQA RLGDSRHRIR KRGLECDGRT  
301 NLCCRQOFFI DFRLIGWNDW IIAPTGYGN YCEGSCPAYL AGVPGSASSF  
351 HTAVVNQYRM RGLNPGTVNS CCIPTKLSTM SMLYFDDEYN IVKRDVPNMI  
401 VEECGCA

## FIG. 24

(SEQ ID NO: 23)

### Human Müllerian Inhibitory Substance (MIS)

1 MRDLPLTSLA LVLSALGALL GTEALRAEEP AVGTSGLIFR EDLDWPPGIP  
51 QEPLCLVALG GDSNGSSSPL RVVGALSAYE QAFLGAVQRA RWGPRDLATF  
101 GVCNTGDRQA ALPSLRRLGA WLRDPGGQRL VVLHLEEVTV EPTPSLRFQE  
151 PPPGGAGPPE LALLVLYPGP GPEVTVTRAG LPGAQSLCPS RDTRYLVLA  
201 DRPAGAWRGS GLALTLPQPG EDSRLSTARL QALLFGDDHR CFTRMTPALL  
251 LLPRSEPAPL PAHGQLDTVP FPPPRPSAEL EESPPSADPF LETLTRLVRA  
301 LRVPPARASA PRLALDPDAL AGFPQGLVNL SDPAALERLL DGEEPLLLLL  
351 RPTAATTGDP APLHDPTSAP WATALARRVA AELQAAAEL RSLPGLPPAT  
401 APLLARLLAL CPGGPGGLGD PLRALLLLKA LQGLRVEWRG RDPRGPGRAQ  
451 RSAGATAADG PCALRELSVD LRAERSVLIP ETYQANNCQG VCGWPQSDRN  
501 PRYGNHVLL LKMQARGAAL ARPPCCVPTA YAGKLLISLS EERISAHVVP  
551 NMVATECGCR

Year	1994	1995	1996	1997	1998	1999	2000	2001	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025	2026	2027	2028	2029	2030	2031	2032	2033	2034	2035	2036	2037	2038	2039	2040	2041	2042	2043	2044	2045	2046	2047	2048	2049	2050	2051	2052	2053	2054	2055	2056	2057	2058	2059	2060	2061	2062	2063	2064	2065	2066	2067	2068	2069	2070	2071	2072	2073	2074	2075	2076	2077	2078	2079	2080	2081	2082	2083	2084	2085	2086	2087	2088	2089	2090	2091	2092	2093	2094	2095	2096	2097	2098	2099	2100
1994	1995	1996	1997	1998	1999	2000	2001	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025	2026	2027	2028	2029	2030	2031	2032	2033	2034	2035	2036	2037	2038	2039	2040	2041	2042	2043	2044	2045	2046	2047	2048	2049	2050	2051	2052	2053	2054	2055	2056	2057	2058	2059	2060	2061	2062	2063	2064	2065	2066	2067	2068	2069	2070	2071	2072	2073	2074	2075	2076	2077	2078	2079	2080	2081	2082	2083	2084	2085	2086	2087	2088	2089	2090	2091	2092	2093	2094	2095	2096	2097	2098	2099	2100	

(SEQ ID NO: 24)

1	MVAGTRCLLA	LLLQVLLGG	AAGLVPELGR	RKFAAAASSGR	PSSQPSDEVL
51	SEFELRLLSM	FGLKQRPTPS	RAVVPYML	DLYRRHSGQP	GSPAPDHRLE
101	RAASRANTVR	SFHHEESLEE	LPETSGKTTR	RFFFNLSSIP	TEEFITSAEL
151	QVFREQMQDA	LGNNSSFHHR	INIYEIIKPA	TANSKFPVTR	LLDTRLVNQN
201	ASRWESFDVT	PAVMRWTAG	HANHGFEV	AHLEEKQGV	KRHVRISRSL
251	HQDEHSWSQI	RPLLVTFGHD	GKGHPLHKRE	KRQAKHKQRK	RLKSSCKRHP
301	<u>LYVDFSDVGW</u>	NDWIVAPPGY	HAFYCHGEC	FPLADHLNST	NHAIVQTLVN
351	SVNSKIPKAC	CVPTELSAIS	MLYLDENEKV	VLKNYQDMVV	EGCGCR

(SEQ ID NO: 25)

1	MAGASRLFL	WLGCFVSLA	QGERPKPPFP	ELRKAVPGDR	TAGGGPDSEL
51	QPQDKVSEHM	LRLYDRYSTV	QAARTPGSLE	GGSQPWRPRL	LREGNTVRSF
101	RAAAAETLER	KGLYIFNLTS	LTKSENILSA	TLYFCIGELG	NISLSCPVS
151	GCSHHAQRKH	IQIDLSAWTL	KFSRNQSPLL	GHLSVDMAKS	HRDIMSWLSK
201	DITQFLRKAK	ENEEFLIGFN	ITSKGRQLPK	RRLPFPEPYI	LVYANDAAIS
251	EPESVSSSLQ	GHRNFPTGTV	PKWDISHIRAA	LSIERRKKRS	TGVLLPLQNN
301	ELPGAQYQYK	KDEVWEERKP	YKTLQAQAPE	KSKNKKKQRK	GPHRKSQTLQ
351	FDEQTLKKAR	RKQWIEPRNC	<u>ARRYLKVDFA</u>	<u>DIGWSEWIS</u>	<u>PKSFDAYYCS</u>
401	GACQFPMPKS	LKPSNHATIO	SIVRAVGVP	GIPECCVPE	<u>KMSSLSILFF</u>
451	<u>DENKNVVLKV</u>	<u>YPNMTVESCA</u>	CR		

13/20

## FIG. 27

(SEQ ID NO: 26)

### Human Bone Morphogenic Protein (BMP)-3b

1 MAHVPARTSP GPGPQLLLLL LPLFLLLLRD VAGSHRAPAW SALPAAADGL  
51 QGDRDLQRHP GDAAATLGPS AQDMVAVMH RLYEKYSRQG ARPGGGNTVR  
101 SFRARLEVVD QKAVYFFNLT SMQDSEMILT ATFFHYSEPP RWPRALEVLC  
151 KPRAKNASGR PLPLGPPTRO HLLFRSLSON TATQGLLRGA MALAPPPRGL  
201 WQAKDISPIV KAARRDGELL LSAQLDSEER DPGVPRPSPY APYILVYAND  
251 LAISEPNSVA VTLQRYDPFP AGDPEPRAAP NNSADPRVRR AAQATGPLQD  
301 NELPGLDERP PRAHAQHFK HQLWPSPFRA LKPRPGRKDR RKKGQEVFMA  
351 ASQVLDFDEK TMQKARRKQW DEPRVCSRRY LKVDFADIGW NEWIISPKSF  
401 DAYYCAGACE FMPKIVRPS NHATIQSIVR AVGIIPGIPE PCCVPDKMNS  
451 LGVLFLENR NVVLKVYPNM SVDTCACR

## FIG. 28

(SEQ ID NO: 27)

### Human Bone Morphogenic Protein (BMP)-4

1 MIPGNRMLMV VLLCQVLLGG ASHASLIPET GKKKVAEIQG HAGGRRSGQS  
51 HELLRDFEAT LLQMFGLERR PQPSKSAVIP DYMRDLYRLQ SGEEEEEQIH  
101 STGLEYPERP ASRANTVRSF HHEEHLENIP GTSENSAFRF LFNLSIPEN  
151 EAISSAELRL FREQVDQGPD WERGFHRINI YEVMKPPAEV VPGHLITRLL  
201 DTRLVHHNVT RWETFDVSPA VLRWTREKQP NYGLAIEVTH LHQTRTHQGQ  
251 HVRISRSLPQ GSGNWAQLRP LLVTFGHDGR GHALTRRRRA KRSPKHHSQR  
301 ARKKKNKNCRR HSLYVDFSDV GWNDWIVAPP GYQAFYCHGD CPFPLADHLN  
351 STNHAIVQTL VNSVNSSIPK ACCVPTLSA ISMLYLDEYD KVVLKQYQEM  
401 VVEGCGCR

14/20

## FIG. 29

(SEQ ID NO: 28)

### Human Bone Morphogenetic Protein (BMP)-5 Precursor

1 MHLTVFLLKG IVGFLWSCWV LVGYAKGGLG DNHVHSSFIY RRLRNHERRE  
51 IQREILSILG LPHRPRPFSP GKQASSAPLF MLDLYNAMTN EENPEESEYS  
101 VRASLAEETR GARKGYASP NGYPRRIQLS RTTPLTTQSP PLASLHDTNF  
151 LNDADMVMSF VNLVERDKDF SHQRRHYKEF RFDLTQIPHG EAVTAAEFRI  
201 YKDRSNNRFE NETIKISIQ IIKEYTNRDA DLFLLDTRKA QALDVGWLVF  
251 DITVTSNHVW INPQNNLGLQ LCAETGDGRS INVKSAGLVG RQGPQSKQPF  
301 MVAFFKASEV LLRSVRAANK RKNQNRNKSS SHQDSSRMSS VGDYNTSEQK  
351 QACKKHELYV SFRDLGWQDW IIAPEGYAAF YCDGECSFPL NAHMNATNHA  
401 IVQTLVHLMF PDHVPKPCCA PTKLNAISVL YFDDSSNVIL KKYRNMVRS  
451 CGCH

## FIG. 30

(SEQ ID NO: 29)

### Human Bone Morphogenetic Protein (BMP)-6/Vgrl

1 SSASDYNSELKTACRKHELYVSFQDLGWqwIIAPKGYAANYCDGECSPP  
51 LNAhtNHAIVQTLVHLMNPEYVPKPCCAPTTKLNAISVLYFDDNSNVikKY  
101 RNMVVRACGCH

21 L1 40  
81 L3

(SEQ ID NO: 30)

1 ANVAENSSSDQRQACKKHELYVSFRDLGWQWIIAPEGYAAYYCEGECAFP  
 21 L1 40  
 51 LNSATNHAIVQTLVHFINPETVPKPCCAPTQLNAISVLYFDDSSNVIKKY  
 81 L3  
 102  
 101 RNMVVRACGCH

(SEQ ID NO: 31)

```

1      MTALPGPLWL  LGLALCALGG  GPGGLRPPPG  CPQRRLGARE  RRDVQREILA

51     VLGLPGRPRP  RAPPAASRLP  ASAPLFMLDL  YHAMAGDDDE  DGAPAERRLG

101    RADLVMSFVN  MVERDRALGH  QEPHWKEFRF  DLTQIPAGEA  VTAAEFRIYK

151    VPSIHLNRT  LHVSMFQVVQ  EQSNRESDLF  FLDLQTLRAG  DEGWLVLDDVT

201    AASDCWLLKR  HKDLGLRLYV  ETEDGHSVDP  GLAGLLGQRA  PRSQQPFFVVT

251    FFRASPSPIR  TPRAVRPLRR  RQPKKSNELP  QANRLPGIFD  DVHGSHGRQV

301    CRRHELYVSF  QDLGWLDWVI  APQGYSAYYC  EGECSFPLDS  CMNATNHAIL

351    QSLVHLMKPN  AVPKACCAPT  KLSATSVLYY  DSSNNVILRK  HRNMVVKACG

401    CH

```

[illegible]

16/20

## FIG. 33

(SEQ ID NO: 32)

### Human Bone Morphogenic Protein (BMP)-10

1 MGSLVLTLCALFCLAAYLVS GSPIMNLEQS PLEEDMSLFG DVFSEQDGVD  
51 FNTLLQSMKD EFLKTLNLS IPTQDSAKVD PPEYMLELYN KFATDRTSMP  
101 SANIIRSFKN EDLFSQPVSF NGLRKYPLLF NVSIPHHEEV IMAELRLYTL  
151 VQRDRMIYDG VDRKITIFEV LESKGDNEGE RNMLVLVS GE IYGTNSEWET  
201 FDVTDAIRRW QKSGSSTHQL EVHIESKHDE AEDASSGRLE IDTSAQNKHN  
251 PLLIVFSD DQ SSDKERKEEL NEMISHEQLP ELDNLGLDSF SSGPGEEALL  
301 QMRSNIIYDS TARIRRNAKG NYCKRTPLYI DFKEIGWDSW IIAPPGYEAY  
351 ECRGVCNYPL AEHLTPTKHA IIQALVHLKN SQKASKACCV PTKLEPISIL  
401 YLDKGVV TYK FKYEGMAVSE CGCR

## FIG. 34

(SEQ ID NO: 33)

### Human Bone Morphogenic Protein (BMP)-11

1 MVLAAPLLLG FLLLALELRP RGEAAEGPAA AAAAAAAAAA AGVGGERS SR  
51 PAPSVAPEPD GCPVCVWRQH SRELRLSEIK SQILSKLRLK EAPNISREVV  
101 KQLLPKAPPL QQILD LHFQ GDALQPEDFL EEDEYHATTE TVISMAQETD  
151 PAVQTDGSPL CCHFHFSPKV MFTKVLKAQL WVYLRPVPRP ATVYLQILRL  
201 KPLTGEGTAG GGGGRRHIR IRSLKIELHS RSGHWQSIDF KQVLHSWFRQ  
251 PQSNWGIEIN AFDPSGTDLA VTSLGPGAEG LHPFMELRVL ENTKRSRRNL  
301 GLDCDEHSSE SRCCRYPLTV DFAFGWDWI IAPKRYKANY CSGQCEYMF M  
351 QKYPHTHLVQ QANPRGSAGP CCTPTKMSPI NMLYFNDKQQ IIYGKIPGMV  
401 VDRCGCS

FIG. 33



17/20

## ***FIG.35***

(SEQ ID NO: 34)

### **I. HUMAN BONE MORPHOGENIC PROTEIN (BMP)-15**

1 MVLLSILRIL FLCELVLFME HRAQMAEGGQ SFIALLAEAP TLPLIEEMLE  
51 ESPGEQPRKP RLLGHSLRYM LELYRRSADS HGHPRENRTI GATMVRLVKP  
101 LTSVARPHRG TWHIQILGFP LRPNRLGYQL VRATVVYRHH LQLTRFNLSC  
151 HVEPWVQKNP TNHFPSSEGD SSKPSLMSNA WKEMDITQLV QQRFWNNKGH  
201 RILRLRFMCQ QQKDSGGLEL WHGTSSLDIA FLLLYFNDTH KSIRKAKFLP  
251 RGMEEFMERE SLLRRTRQAD GISA EVTASS SKHSGPENNQ CSLHPFQISE  
301 RQLGWDHWII APPFYTPNYC KGTCLRVL RD GLNSPNHAI QNLINQLVDQ  
351 SVPRPSCVPY KYVPISVLMI EANGSILYKE YEGMIAESCT CR

## ***FIG.36***

(SEQ ID NO: 35)

### **Human Norrie Disease Protein (NDP)**

**[Norrin]**

1 MRKHVLAASF SMLSLLVIMG DTDSKTDSSF IMDSDP RR CM RHHYVDSISH  
51 PLYKCSSKMV LLARCEGHCS QASRSEPLVS FSTVLKQPFR SSCHCCRPQT  
101 SKLKALRLRC SGGMRLTATY RYILSCHCEE CNS

18/20

## FIG. 37

(SEQ ID NO: 36)

### Human Growth Differentiation Factor (GDF)-1

1 MPPPQQGPCG HLLLLLLALL LPSLPLTRAP VPPGPAAALL QALGLRDEPQ  
51 GAPRLRPVPP VMWRLFRRRD PQETRSGSRR TSPGVTLQPC HVEELGVAGN  
101 IVRHIPDRGA PTRASEPVSA AGHCPewTVV FDLSAVEPAE RPSRARLELR  
151 FAAAAAAPE GGWELSVAQA GQGAGADPGP VLLRQLVPAL GPPVRAELLG  
201 AAWARNASWP RSLRLALALR PRAPAACARL AEASLLLVTL DPRLCHPLAR  
251 PRRDAEPVLG GPGGACRAR RLYVSFREV G WHRWVIAPRG FLANYCQGQC  
301 ALPVALSGSG GPPALNHA VL RALMHAAAPG AADLPCCVPA RLSPISVLFF  
351 DNSDNVVLRO YEDMVVDECG CR

## FIG. 38

(SEQ ID NO: 37)

### Human Growth Differentiation Factor (GDF)-5 Precursor

1 MRLPKLLTFL LWYLA WLDLE FICTVLGAPD LGQRPQGS RP GLAKAEAKER  
51 PPLARNVFRP GGHSYGGGAT NANARAKGGT GQTGGTLQPK KDEPKKLPPR  
101 PGGPEPKPGH PPQTRQATAR TVTPKGQLPG GKAPPKAGSV PSSFLLKKAR  
151 EPGPPREPKE PFRPPPITPH EYMLSLYRTL SDADRKGGNS SVKLEAGLAN  
201 TITSFIDKGQ DDRGPVVRKQ RYVFDISALE KDGLLGAE LR ILRKKPSDTA  
251 KPAVPRSRR AQLKLSSCPS GRQPAALLDV RSVPGLDGSG WEVFDIWKLF  
301 RNFKN SAQLC LELEAWERGR TVDLRGLGFD RAARQVHEKA LFLVFGRTKK  
351 RDLFFNEIKA RSGQDDKTVY EYLFSQRRKR RAPSATRQ GK RPSKNL KARC  
401 SRKALHVNFK DMGWDDWIIA PLEYEAFHCE GLCEFPLRSH LEPTNHA VIQ  
451 TLMNSMDPES TPPTCCVPTR LSPISILFID SANNVVKQY EDMVVESCGC  
501 R

19/20

## FIG. 39

(SEQ ID NO: 38)

### Human Growth Differentiation Factor (GDF)-8 [Myostatin]

1 MQKLQLCVYI YLFMLIVAGP VDLNENSEQK ENVEKEGLCN ACTWRQNTKS  
51 SRIEAIKIQI LSKLRLETAP NISKDVIRQL LPKAPPLREL IDQYDVQRDD  
101 SSDGSLEDDD YHATTETIIT MPTESDFLMQ VDGKPKCCFF KFSSKIQYNK  
151 VVKAQLWIYL RPYETPTTVF VQILRLIKPM KDGTRYTGIR SLKLDMPNGT  
201 GIWQSIDVKT VLQNLWKQPE SNLGIEIKAL DENGHD LAVT FPGPGEDGLN  
251 PFLEVKVTD T PKRSRRDFGL DCDEHSTESR CCRYPLTVDF EAFGWDWIIA  
301 PKRYKANYCS GECEVFVLQK YPHTHLVHQA NPRGSAGPCC TPTKMSPINM  
351 LYFNGKEQII YGKIPAMVVD RCGCS

## FIG. 40

(SEQ ID NO: 39)

### Human Growth Differentiation Factor (GDF)-9

1 MARPNKFLW FCCFAWLCFP ISLGSQASGG EAQIAASAEL ESGAMPWSLL  
51 QHIDERDRAG LLPALFKVLS VGRGGSPRLQ PDSRALHYMK KLYKTYATKE  
101 GIPKSNRSHL YNTVRLFTPC TRHKQAPGDQ VTGILPSVEL LFNLDRIITV  
151 EHLLKSVLLY NINNSVSFSS AVKVCVNLMI KEPKSSSRTL GRAPYSFTFN  
201 SQFEFGKKHK WIQIDVTSLL QPLVASNKRS IHMSINF TCM KDQLEHPSAQ  
251 NGLFNMTLVS PSLILYLNDT SAQAYHSWYS LHYKRRPSQG PDQERSLSAY  
301 PVGEEAAEDG RSSHHRHRRG QETVSSELKK PLGPASFNLS EYFRQFLLPQ  
351 NECELHDFRL SFSQLKWDNW IVAPHRYNPR YCKGDCPRAV GHRYGSPVHT  
401 MVQNIIEYKL DSSVPRPSCV PAKYSPLSVL TIEPDGSIAY KEYEDMIATK  
451 CTRC

20/20

## FIG. 41

(SEQ ID NO: 40)

### Human Artemin (GDNF)

1 MPGLISARGQ PLLEVLPPQA HLGALFLPEA PLGLSAQPAL WPTLAALALL  
51 SSVAEASLGS APRSPAPREG PPPVLASPAG HLPGGRTARW CSGRRARRPPP  
101 QPSRPAPPPP APPSALPRGG RAARAGGPGS RARAAGARGC RLRSOLVPVR  
151 ALGLGHRSD LVRFRFCSGS CRRARSPHDL SLASLLGAGA LRPPPGSRPV  
201 SQPCCRPTRY EAVSFMDVNS TWRTVDRLSA TACGCLG

## FIG. 42

(SEQ ID NO: 41)

### Human Glial Cell Derived Factor (GDNF)

[Persephin]

1 MAVGKFLGGS LLLLSLQLGQ GWGPDARGVP VADGEFSSEQ VAKAGGTWLG  
51 THRPLARLRR ALSGPCQLWS LTLSVAELGL GYASEEKVIF RYCAGSCPRG  
101 ARTQHGLALA RLQGQGRAHG GPCCRPTRYT DVAFLDDRHR WORLPOLSAA  
151 ACGCGG

FIG. 41 "SEQ ID NO: 40"